

A. MARSCHEL
RAW SEQUENCE LISTING
PATENT APPLICATION US/08/869,275renew
DATE: 04/18/2000
TIME: 09:28:06

INPUT SET: S35320.raw

This Raw Listing contains the General
Information Section and up to the first 5 pages.

SEQUENCE LISTING

ENTERED

(1) General Information:

(i) APPLICANT: Carl T. Wittwer, Kirk M. Ririe, Randy P.
Rasmussen, and David R. Hillyard(ii) TITLE OF INVENTION: SYSTEM AND METHOD FOR CARRYING OUT
AND MONITORING BIOLOGICAL PROCESSES

(iii) NUMBER OF SEQUENCES: 6

(iv) CORRESPONDENCE ADDRESS:

(A) ADDRESSEE: Thorpe, North & Western
(B) STREET: 9035 South 700 East, Suite 200
(C) CITY: Sandy
(D) STATE: Utah
(E) COUNTRY: USA
(F) ZIP: 84070

(v) COMPUTER READABLE FORM:

(A) MEDIUM TYPE: Diskette, 3.5 inch, 1.4 Mb storage
(B) COMPUTER: Toshiba 200CDS
(C) OPERATING SYSTEM: DOS 7.0
(D) SOFTWARE: Word Perfect 7.0

(vi) CURRENT APPLICATION DATA:

(A) APPLICATION NUMBER:
(B) FILING DATE: June 4, 1997
(C) CLASSIFICATION:

(vii) PRIOR APPLICATION DATA:

(A) APPLICATION NUMBER:
(B) FILING DATE:

(viii) ATTORNEY/AGENT INFORMATION:

(A) NAME: Grant R. Clayton
(B) REGISTRATION NUMBER: 32,462
(C) REFERENCE/DOCKET NUMBER: 8616CIP5

(ix) TELECOMMUNICATION INFORMATION:

(A) TELEPHONE: (801)566-6633
(B) TELEFAX: (801)566-0750

(2) INFORMATION FOR SEQ ID NO:1:

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47 (i) SEQUENCE CHARACTERISTICS:
48 (A) LENGTH: 20
49 (B) TYPE: nucleic acid
50 (C) STRANDEDNESS: single-stranded
51 (D) TOPOLOGY: linear
52
53 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:
54
55 CGTGGTGGAC TTCTCTCAAT 20
56
57 (2) INFORMATION FOR SEQ ID NO:2:
58
59 (i) SEQUENCE CHARACTERISTICS:
60 (A) LENGTH: 20
61 (B) TYPE: nucleic acid
62 (C) STRANDEDNESS: single-stranded
63 (D) TOPOLOGY: linear
64
65 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:2:
66
67 AGAAGATGAG GCATAGCAGC 20
68
69
70 (2) INFORMATION FOR SEQ ID NO:3:
71
72 (i) SEQUENCE CHARACTERISTICS:
73 (A) LENGTH: 35
74 (B) TYPE: nucleic acid
75 (C) STRANDEDNESS: single-stranded
76 (D) TOPOLOGY: linear
77
78 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:3:
79
80 CAAACAGACA CCATGGTGCA CCTGACTCCT GAGGA 35
81
82
83 (2) INFORMATION FOR SEQ ID NO:4:
84
85 (i) SEQUENCE CHARACTERISTICS:
86 (A) LENGTH: 30
87 (B) TYPE: nucleic acid
88 (C) STRANDEDNESS: single-stranded
89 (D) TOPOLOGY: linear
90
91 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:4:
92
93 AAGTCTGCCG TTACTGCCCT GTGGGGCAAG 30
94
95
96
97 (2) INFORMATION FOR SEQ ID NO:5:
98
99 (i) SEQUENCE CHARACTERISTICS:

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100 (A) LENGTH: 22
101 (B) TYPE: nucleic acid
102 (C) STRANDEDNESS: single-stranded
103 (D) TOPOLOGY: linear
104
105 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:5:
106
107 GGTTGGCCAA TCTACTCCCA GG 22
108
109
110 (2) INFORMATION FOR SEQ ID NO:6:
111
112 (i) SEQUENCE CHARACTERISTICS:
113 (A) LENGTH: 20
114 (B) TYPE: nucleic acid
115 (C) STRANDEDNESS: single-stranded
116 (D) TOPOLOGY: linear
117
118 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:6:
119
120 GCTCACTCAG TGTGGCAAAG 20
121
122

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SEQUENCE VERIFICATION REPORT
PATENT APPLICATION US/08/869,275

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Line

Error

Original Text